

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 09/976,935A  
Source: IFW/6  
Date Processed by STIC: 3-30-05

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IFW16

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/976,935A**

DATE: 03/30/2005  
TIME: 15:14:26

Input Set : A:\36470Asubseq.txt  
Output Set: N:\CRF4\03302005\I976935A.raw

5 <110> APPLICANT: Staunton, Donald E.  
9 <120> TITLE OF INVENTION: MATERIALS AND METHODS TO MODULATE LIGAND BINDING/ENZYMATIC

## ACTIVITY

10 OF ALPHA/BETA PROTEINS CONTAINING AN ALLOSTERIC REGULATORY SITE  
14 <130> FILE REFERENCE: 27866/36470A  
18 <140> CURRENT APPLICATION NUMBER: 09/976,935A  
20 <141> CURRENT FILING DATE: 2001-10-12  
24 <150> PRIOR APPLICATION NUMBER: US 60/239,750  
26 <151> PRIOR FILING DATE: 2000-10-12  
30 <160> NUMBER OF SEQ ID NOS: 36  
34 <170> SOFTWARE: PatentIn version 3.1  
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72 <213> ORGANISM: Q327A  
76 <400> SEQUENCE: 3  
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84 <212> TYPE: DNA  
86 <213> ORGANISM: I332A  
90 <400> SEQUENCE: 4  
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94 <210> SEQ ID NO: 5  
96 <211> LENGTH: 32  
98 <212> TYPE: DNA  
100 <213> ORGANISM: F333A  
104 <400> SEQUENCE: 5  
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108 <210> SEQ ID NO: 6  
110 <211> LENGTH: 33  
112 <212> TYPE: DNA

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128 <213> ORGANISM: Primer
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136 <210> SEQ ID NO: 8
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140 <212> TYPE: DNA
142 <213> ORGANISM: Primer
146 <400> SEQUENCE: 8
147 aatttctcgaa ggtctccaac cgtgccttcc 30
150 <210> SEQ ID NO: 9
152 <211> LENGTH: 27
154 <212> TYPE: PRT
156 <213> ORGANISM: Amino acid insertion
160 <400> SEQUENCE: 9
162 Pro Lys Gly Arg His Arg Gly Val Thr Val Val Arg Ser His His Gly
163 1 5 10 15
166 Val Leu Ile Cys Ile Gln Val Leu Val Arg Arg
167 20 25
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176 <213> ORGANISM: primer Eo26-H3
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181 gaggggaagc tttagtgggcc 20
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190 <213> ORGANISM: primer Eo-24
194 <400> SEQUENCE: 11
195 gaagttggcc tgagcctgg 19
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200 <211> LENGTH: 25
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204 <213> ORGANISM: E-cad 5' #1
208 <400> SEQUENCE: 12
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214 <211> LENGTH: 27
216 <212> TYPE: DNA
218 <213> ORGANISM: E-cad 3' #1
222 <400> SEQUENCE: 13
223 ctgcacatgg tctggccgc ctctctc 27
226 <210> SEQ ID NO: 14

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228 <211> LENGTH: 45  
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 246 <213> ORGANISM: Primer Ecad3' (Xho)  
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 251 aggcgctcga gaatccccag aatggcagga att 33  
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 260 <213> ORGANISM: primer MAdCAM-1 5'#1  
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 306 <400> SEQUENCE: 19  
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 312 <211> LENGTH: 28  
 314 <212> TYPE: DNA  
 316 <213> ORGANISM: Primer Alpha1.5  
 320 <400> SEQUENCE: 20  
 321 gactttcagc ggcccggtgg aagacatg 28  
 324 <210> SEQ ID NO: 21  
 326 <211> LENGTH: 28  
 328 <212> TYPE: DNA  
 330 <213> ORGANISM: Primer Alpha1.3  
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 338 <210> SEQ ID NO: 22  
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Input Set : A:\36470Asubseq.txt

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354 <211> LENGTH: 33	
356 <212> TYPE: DNA	
358 <213> ORGANISM: A1.3Bam	
362 <400> SEQUENCE: 23	
363 atatggatcc ctaagacatt tccatttcaa atg	33
366 <210> SEQ ID NO: 24	
368 <211> LENGTH: 29	
370 <212> TYPE: DNA	
372 <213> ORGANISM: A2.5Nde	
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382 <211> LENGTH: 32	
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390 <400> SEQUENCE: 25	
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404 <400> SEQUENCE: 26	
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410 <211> LENGTH: 31	
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422 <210> SEQ ID NO: 28	
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426 <212> TYPE: DNA	
428 <213> ORGANISM: Primer specific for 5'EchHPPK	
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433 gtagatgaca gtggcgtata tt	22
436 <210> SEQ ID NO: 29	
438 <211> LENGTH: 23	
440 <212> TYPE: DNA	
442 <213> ORGANISM: Primer specific for 3'EchHPPK	
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Input Set : A:\36470Asubseq.txt  
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456 <213> ORGANISM: amino acid sequence of ???  
 460 <400> SEQUENCE: 30  
 462 Met Thr Val Ala Tyr Ile Ala Ile Gly Ser Asn Leu Ala Ser Pro Leu  
 463 1 5 10 15  
 466 Glu Gln Val Asn Ala Ala Leu Lys Ala Leu Gly Asp Ile Pro Glu Ser  
 467 20 25 30  
 470 His Ile Leu Thr Val Ser Ser Phe Tyr Arg Thr Pro Pro Leu Gly Pro  
 471 35 40 45  
 474 Gln Asp Gln Pro Asp Tyr Leu Asn Ala Ala Val Ala Leu Glu Thr Ser  
 475 50 55 60  
 478 Leu Ala Pro Glu Glu Leu Leu Asn His Thr Gln Arg Ile Glu Leu Gln  
 479 65 70 75 80  
 482 Gln Gly Arg Val Arg Lys Ala Glu Arg Trp Gly Pro Arg Thr Leu Asp  
 483 85 90 95  
 486 Leu Asp Ile Met Leu Phe Gly Asn Glu Val Ile Asn Thr Glu Arg Leu  
 487 100 105 110  
 490 Thr Val Pro His Tyr Asp Met Lys Asn Arg Gly Phe Met Leu Trp Pro  
 491 115 120 125  
 494 Leu Phe Glu Ile Ala Pro Glu Leu Val Phe Pro Asp Gly Glu Met Leu  
 495 130 135 140  
 498 Arg Gln Ile Leu His Thr Arg Ala Phe Asp Lys Leu Asn Lys Trp  
 499 145 150 155  
 502 <210> SEQ ID NO: 31  
 504 <211> LENGTH: 10  
 506 <212> TYPE: PRT  
 508 <213> ORGANISM: ????  
 512 <400> SEQUENCE: 31  
 514 Met Gly His His His His His Gly Gly  
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 518 <210> SEQ ID NO: 32  
 520 <211> LENGTH: 52  
 522 <212> TYPE: DNA  
 524 <213> ORGANISM: 5'EcHisHPPK  
 528 <400> SEQUENCE: 32  
 529 cgccatggc caccaccacc accaccacgg cgccatgaca gtggcgata tt 52  
 532 <210> SEQ ID NO: 33  
 534 <211> LENGTH: 29  
 536 <212> TYPE: DNA  
 538 <213> ORGANISM: 3'EcXhoHPPK  
 542 <400> SEQUENCE: 33  
 543 cggctcgagt taccatttgt ttaatttgt 29  
 546 <210> SEQ ID NO: 34  
 548 <211> LENGTH: 169  
 550 <212> TYPE: PRT  
 552 <213> ORGANISM: amino acid sequence of His(6)-HPPK gene  
 556 <400> SEQUENCE: 34  
 558 Met Gly His His His His His Gly Gly Met Thr Val Ala Tyr Ile  
 559 1 5 10 15  
 562 Ala Ile Gly Ser Asn Leu Ala Ser Pro Leu Glu Gln Val Asn Ala Ala

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